

09/763957

JC03 Rec'd PCT/PTO 28 FEB 2001
PCT/AU99/00705

WO 00/12714

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SEQUENCE LISTING

<110> THE UNIVERSITY OF QUEENSLAND

<120> A NOVEL PLANT PROMOTER AND USES THEREFOR

<130> 2209376/EJR

<140>

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<150> PP5572

<151> 1998-08-31

<160> 9

<170> PatentIn Ver. 2.0

<210> 1

<211> 1923

<212> DNA

<213> Plant

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<221> CDS

<222> (88)..(1539)

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Pro Leu Leu Ser Lys Met Ala Ile Gly Asp Gly His Gly Glu Ser Ser
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cca tac ttt gat gga tgg aag gct tat gat caa aac ccc ttt cat ccc 210
Pro Tyr Phe Asp Gly Trp Lys Ala Tyr Asp Gln Asn Pro Phe His Pro
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Thr Asp Asn Pro Asn Gly Val Met Gln Met Gly Leu Ala Glu Asn Gln
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 tcc att tgc act cca gaa gga ata aat gat ttc agg gcc ata gct aac	354
Ser Ile Cys Thr Pro Glu Gly Ile Asn Asp Phe Arg Ala Ile Ala Asn	
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 ttt cag gat tat cat ggt ctg gcc gag ttc aga aat gct gtg gct aaa	402
Phe Gln Asp Tyr His Gly Leu Ala Glu Phe Arg Asn Ala Val Ala Lys	
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Phe Met Ala Arg Thr Arg Gly Asn Arg Ile Thr Phe Asp Pro Asp Arg	
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Ile Val Met Ser Gly Gly Ala Thr Gly Ala His Glu Val Thr Ala Phe	
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Cys Leu Ala Asp Pro Gly Glu Ala Phe Leu Val Pro Ile Pro Tyr Tyr	
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Pro Gly Phe Asp Arg Asp Leu Arg Trp Arg Thr Gly Val Lys Leu Val	
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Pro Val Met Cys Asp Ser Ser Asn Asn Phe Val Leu Thr Lys Glu Ala	
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Leu Glu Asp Ala Tyr Glu Lys Ala Arg Glu Asp Asn Ile Arg Val Lys	
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 ggc tta ctg atc acc aat cca tca aat cca tta ggc aca atc atg gac	738
Gly Leu Leu Ile Thr Asn Pro Ser Asn Pro Leu Gly Thr Ile Met Asp	
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 aga aag aca ctg aga acc gtc gtc agc ttc atc aat gag aag cgt atc	786
Arg Lys Thr Leu Arg Thr Val Val Ser Phe Ile Asn Glu Lys Arg Ile	
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225	230
 cac ctt gta tgt gat gaa ata tat gct gca aca gtt ttc agc caa ccc	834
His Leu Val Cys Asp Glu Ile Tyr Ala Ala Thr Val Phe Ser Gln Pro	

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cca act ttc gac tct gaa acg gag ctt tgg aaa gtt atc att cat gaa Pro Thr Phe Asp Ser Glu Thr Glu Leu Trp Lys Val Ile Ile His Glu 380	385	390	1266	
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 Leu Lys Thr Arg Arg Phe Asp Asp Ile Thr Met Ser Pro His Ser Pro
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Asn Arg Ile Thr Phe Asp Pro Asp Arg Ile Val Met Ser Gly Gly Ala			
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Thr Gly Ala His Glu Val Thr Ala Phe Cys Leu Ala Asp Pro Gly Glu			
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Ala Phe Leu Val Pro Ile Pro Tyr Tyr Pro Gly Phe Asp Arg Asp Leu			
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Asn Asn Phe Val Leu Thr Lys Glu Ala Leu Glu Asp Ala Tyr Glu Lys			
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Ser Asn Pro Leu Gly Thr Ile Met Asp Arg Lys Thr Leu Arg Thr Val			
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Val Ser Phe Ile Asn Glu Lys Arg Ile His Leu Val Cys Asp Glu Ile			
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